

05/10  
0606  
#5



OIPE

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/032,256A

DATE: 06/03/2002

TIME: 13:22:15

Input Set : A:\M2335hnl.app

Output Set: N:\CRF3\06032002\J032256A.raw

ENTERED

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3 <110> APPLICANT: CHODOSH, Lewis A
4   GARDNER, Heather P
6 <120> TITLE OF INVENTION: HORMONALLY UP-REGULATED, NEU-TUMOR-ASSOCIATED KINASE
8 <130> FILE REFERENCE: 22253-70421
10 <140> CURRENT APPLICATION NUMBER: 10/032,256A
C--> 11 <141> CURRENT FILING DATE: 2002-05-21
13 <150> PRIOR APPLICATION NUMBER: 60/257,073
14 <151> PRIOR FILING DATE: 2000-12-21
16 <160> NUMBER OF SEQ ID NOS: 18
18 <170> SOFTWARE: PatentIn Ver. 2.1
20 <210> SEQ ID NO: 1
21 <211> LENGTH: 5024
22 <212> TYPE: DNA
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28 ggggcgatgg aggcgcggag gacacgacca ggccggcggc ggccctgcgag ggaagtctcc 180
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57 ggacactctc ccaggggctg ctgtccggaa gtacctcacc tctccaaact ccaactgcatt 1920
58 ccagctggt ctcttttgcc cacgaagaaa agaacagccc cccgaaagag gaggggtgtgt 1980
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115 <213> ORGANISM: Murinae gen. sp.
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119 1 5 10 15
121 Gly Gly Asp Gly Gly Ala Glu Asp Thr Thr Arg Pro Ala Ala Ala Cys
122 20 25 30
124 Glu Gly Ser Phe Leu Pro Ala Trp Val Ser Gly Val Ser Arg Glu Arg
125 35 40 45
127 Leu Arg Asp Phe Gln His His Lys Arg Val Gly Asn Tyr Leu Ile Gly
128 50 55 60
130 Ser Arg Lys Leu Gly Glu Gly Ser Phe Ala Lys Val Arg Glu Gly Leu
131 65 70 75 80
133 His Val Leu Thr Gly Glu Lys Val Ala Ile Lys Val Ile Asp Lys Lys
134 85 90 95
136 Arg Ala Lys Lys Asp Thr Tyr Val Thr Lys Asn Leu Arg Arg Glu Gly
137 100 105 110
139 Gln Ile Gln Gln Met Ile Arg His Pro Asn Ile Thr Gln Leu Leu Asp
140 115 120 125
142 Ile Leu Glu Thr Glu Asn Ser Tyr Tyr Leu Val Met Glu Leu Cys Pro
143 130 135 140
145 Gly Gly Asn Leu Met His Lys Ile Tyr Glu Lys Lys Arg Leu Asp Glu
146 145 150 155 160
148 Ala Glu Ala Arg Arg Tyr Ile Arg Gln Leu Ile Ser Ala Val Glu His
149 165 170 175
151 Leu His Arg Ala Gly Val Val His Arg Asp Leu Lys Ile Glu Asn Leu
152 180 185 190
154 Leu Leu Asp Glu Asp Asn Asn Ile Lys Leu Ile Asp Phe Gly Leu Ser
155 195 200 205
157 Asn Cys Ala Gly Ile Leu Gly Tyr Ser Asp Pro Phe Ser Thr Gln Cys
158 210 215 220
160 Gly Ser Pro Ala Tyr Ala Ala Pro Glu Leu Leu Ala Arg Lys Lys Tyr
161 225 230 235 240
163 Gly Pro Lys Ile Asp Val Trp Ser Ile Gly Val Asn Met Tyr Ala Met
164 245 250 255
166 Leu Thr Gly Thr Leu Pro Phe Thr Val Glu Pro Phe Ser Leu Arg Ala
167 260 265 270
169 Leu Tyr Gln Lys Met Val Asp Lys Ala Met Asn Pro Leu Pro Thr Gln
170 275 280 285
172 Leu Ser Thr Gly Ala Val Asn Phe Leu Arg Ser Leu Leu Glu Pro Asp

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## RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/032,256A

DATE: 06/03/2002

TIME: 13:22:16

Input Set : A:\M2335hn1.app

Output Set: N:\CRF3\06032002\J032256A.raw

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176 305      310      315      320
178 Asn Glu Asn Tyr Thr Gly Lys Val Pro Cys Asn Val Thr Tyr Pro Asn
179      325      330      335
181 Arg Ile Ser Leu Glu Asp Leu Ser Pro Ser Val Val Leu His Met Thr
182      340      345      350
184 Glu Lys Leu Gly Tyr Lys Asn Ser Asp Val Ile Asn Thr Val Leu Ser
185      355      360      365
187 Asn Arg Ala Cys His Ile Leu Ala Ile Tyr Phe Leu Leu Asn Lys Lys
188      370      375      380
190 Leu Glu Arg Tyr Leu Ser Gly Lys Ser Asp Ile Gln Asp Ser Ile Cys
191 385      390      395      400
193 Tyr Lys Thr Gln Leu Tyr Gln Ile Glu Lys Cys Arg Ala Thr Lys Glu
194      405      410      415
196 Pro Tyr Glu Ala Ser Leu Asp Thr Trp Thr Arg Asp Phe Glu Phe His
197      420      425      430
199 Ala Val Gln Asp Lys Lys Pro Lys Glu Gln Glu Lys Arg Gly Asp Phe
200      435      440      445
202 Leu His Arg Pro Phe Ser Lys Lys Leu Asp Lys Asn Leu Pro Ser His
203      450      455      460
205 Lys Gln Pro Ser Pro Ser Leu Ile Thr Gln Leu Gln Ser Thr Lys Ala
206 465      470      475      480
208 Leu Leu Lys Asp Arg Lys Ala Ser Lys Ser Gly Phe Pro Asp Lys Asp
209      485      490      495
211 Ser Phe Val Cys Arg Asn Leu Phe Arg Lys Thr Ser Asp Ser Asn Cys
212      500      505      510
214 Val Ala Ser Ser Ser Met Glu Phe Ile Pro Val Pro Pro Arg Thr
215      515      520      525
217 Pro Arg Ile Val Lys Lys Leu Glu Pro His Gln Pro Gly Pro Gly Ser
218      530      535      540
220 Ala Ser Ile Leu Pro Lys Glu Glu Pro Leu Leu Asp Met Val Arg
221 545      550      555      560
223 Ser Phe Glu Ser Val Asp Arg Glu Asp His Ile Glu Leu Leu Ser Pro
224      565      570      575
226 Ser His His Tyr Arg Ile Leu Ser Ser Pro Val Ser Leu Ala Arg Arg
227      580      585      590
229 Asn Ser Ser Glu Arg Thr Leu Ser Gln Gly Leu Leu Ser Gly Ser Thr
230      595      600      605
232 Ser Pro Leu Gln Thr Pro Leu His Ser Thr Leu Val Ser Phe Ala His
233      610      615      620
235 Glu Glu Lys Asn Ser Pro Pro Lys Glu Glu Gly Val Cys Ser Pro Pro
236 625      630      635      640
238 Pro Val Pro Ser Asn Gly Leu Leu Gln Pro Leu Gly Ser Pro Asn Cys
239      645      650      655
241 Val Lys Ser Arg Gly Arg Phe Pro Met Met Gly Ile Gly Gln Met Leu
242      660      665      670
244 Arg Lys Arg His Gln Ser Leu Gln Pro Ser Ser Glu Arg Ser Leu Asp
245      675      680      685

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Output Set: N:\CRF3\06032002\J032256A.raw

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255 <211> LENGTH: 10
256 <212> TYPE: RNA
257 <213> ORGANISM: Unknown Organism
259 <220> FEATURE:
260 <223> OTHER INFORMATION: Description of Unknown Organism:Kozak consensus
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268 <211> LENGTH: 6
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270 <213> ORGANISM: Unknown Organism
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273 <223> OTHER INFORMATION: Description of Unknown Organism:polyadenylation
274      signal
276 <400> SEQUENCE: 4
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281 <211> LENGTH: 6
282 <212> TYPE: DNA
283 <213> ORGANISM: Murinae gen. sp.
285 <400> SEQUENCE: 5
286 aataca                      6
289 <210> SEQ ID NO: 6
290 <211> LENGTH: 6
291 <212> TYPE: PRT
292 <213> ORGANISM: murine Hunk;fragment
294 <400> SEQUENCE: 6
295 Asp Leu Lys Pro Glu Asn
296      1                      5
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300 <211> LENGTH: 21
301 <212> TYPE: DNA
302 <213> ORGANISM: Artificial Sequence
304 <220> FEATURE:
305 <221> NAME/KEY: misc_feature
306 <222> LOCATION: (17)
307 <223> OTHER INFORMATION: n is a, c, g, or t
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310 <223> OTHER INFORMATION: Description of Artificial Sequence:degenerate
311      oligonucleotide primer PTKIa
313 <400> SEQUENCE: 7
W--> 314 gggcccggat ccacmgngay y                      21
317 <210> SEQ ID NO: 8

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RAW SEQUENCE LISTING ERROR SUMMARY  
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Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:7; N Pos. 17